

FIGURE 1

1 GAATTCAAGA CCAGCCTGGA CAACTTGGAA GAACCCGGTC TCTACAAAAA ATACAAAATT
 61 AGCTGGGATT GGGTGCGGTG GCTCATGCCT ATAATCCCAG CACTTTGGGA GCCTGAGGTG
 121 GGTGGATCAC CTGAAGTCAG GAGTTCAGA CTAGCCTGGC CAACATGGTG AAACCCCTATC
 181 TCTACTGAAA ATACAAAAAG CTAGACGTGG TGGCACACAC CTGTAATCCC AGCTACTTAG
 241 GAGGCTGAGG CAGGAGAATT GCTTGAAGCC TAGAGGTGAA GGTGTAGTG AGCCGAGATT
 301 GCATCATTCG ACAAATGGAGG GGAGCCACCA GCCTGGGCAA CAAGAGGAAA TCTCCGTCTC
 361 CAAAAA AAAA AAAA AAAGAATTAG GCTGGGTGGT GCCTGTAGTC CCAGCTACTT
 421 GGGAGGCAGG GGGTCCACTT GATGTCGAGA CTGCAGTGAG CCATGATCCT GCCACTGCAC
 481 TCCGGCCTGG GCAACAGAGT GAGACCCTGT CTAAAGAAAA AAAAAATAAA GCAACATATC
 541 CTGAACAAAG GATCCTCCAT AACGTTCCCA CCAGATTCTT AATCAGAAAC ATGGAGGCCA
 601 GAAAGCAGTG GAGGAGGACG ACCCTCAGCG AGCCCGGGAG GATGTTGTCA CAGGCTGGGG
 661 CAAGGGCCTT CCGCTACCA ACTGGGAGCT CTGGGAACAG CCCTGTTGCA AACAAGAAGC
 721 CATAGCCCGG CCAGAGCCCA GGAATGTGGG CTGGGCTGGG AGCAGCCTCT GGACAGGAGT
 781 GGTCCCATCC AGGAAACCTC CGGCATGGCT GGGAAAGTGG GTACTTGGTG CCGGGTCTGT
 841 ATGTGTGTGT GACTGGTGTG TGTGAGAGAG AATGTGTGCC CTAAGTGTCA GTGTGAGTCT
 901 GTGTATGTGT GAATATTGTC TTTGTGTGGG TGATTTTCTG CGTGTGTAAT CCGTGTCCCTG
 961 CAAGTGTGAA CAAGTGGACA AGTGTCTGGG AGTGGACAAG AGATCTGTGC ACCATCAGGT
 1021 GTGTGCATAG CGTCTGTGCA TGTCAAGAGT GCAAGGTGAA GTGAAGGGAC CAGGCCCATG
 1081 ATGCCACTCA TCATCAGGAG CTCTAAGGCC CCAGGTAAGT GCCAGTGACA GATAAGGGTG
 1141 CTGAAGGTCA CTCTGAGAGT GGCAGGTGGG GGTAGGGAAA GGGCAAGGCC ATGTTCTGGA
 1201 GGAGGGGTTG TGACTACATT AGGGTGTATG AGCCTAGCTG GGAGGTGGAT AGCCGGGTCC
 1261 ACTGAAACCC TGGTTATCCC AGAAGGCTTT GCAGGCTTCA GGAGCTTGA GTGGGGAGAG
 1321 GGGGTGACTT CTCCGACCAG GCCCTCCAC CGGCCTACCC TGGGTAAGGG CCTGGAGCAG
 1381 GAAGCAGGGG CAAGAACCTC TGGAGCAGCC CATAACCGCC CTGGCCTGAC TCTGCCACTG
 1441 GCAGCACAGT CAACACAGCA GGTTCATCA CAGCAGAGGG CAAAGGCCAT CATCAGCTCC
 1501 CTTTATAAGG GAAGGGTCAC GCGCTCGGTG TGCTGAGAGT GTCTGCCTG GTCCTCTGTG
 1561 CCTGGTGGGG TGGGGGTGCC AGGTGTGTCC AGAGGAGCCC ATTTGGTAGT GAGGCAGGTA
 1621 TGGGGCTAGA AGCACTGGTG CCCCTGGCCG TGATAGTGGC CATCTTCCTG CTCCTGGTGG
 1681 ACCTGATGCA CCGGCGCCAA CGCTGGGCTG CACGCTACCC ACCAGGCCCC CTGCCACTGC
 1741 CCGGGCTGGG CAACCTGCTG CATGTGGAGT TCCAGAACAC ACCATACTGC TTCGACCAGG
 1801 TGAGGGAGGA GGTCTTGGAG GGCGGCAGAG GTGCTGAGGC TCCCCTACCA GAAGCAAACA
 1861 TGGATGGTGG GTGAAACCAC AGGCTGGACC AGAAGCCAGG CTGAGAAGGG GAAGCAGGTT
 1921 TGGGGGACGT CCTGGAGAAG GGCATTTATA CATGGCATGA AGGACTGGAT TTTCCAAAGG
 1981 CCAAGGAAGA TAGGGCAAG GGCCTGGAGG TGGAGCTGGA CTTGGCAGTG GGCATGCAAG
 2041 CCCATTGGGC AACATATGTT ATGGAGTACA AAGTCCCCTC TGCTGACACC AGAAGGAAAG
 2101 GCCTTGGGAA TGGAAGATGA GTTAGTCCTG AGTGCCGTTT AAATCACGAA ATCGAGGATG
 2161 AAGGGGGTGC AGTGACCCGG TTCAAACCTT TTGCACTGTG GGTCTCGGG CCTCACTGCC
 2221 TCACCGGCAT GGACCATCAT CTGGGAATGG GATGCTAACT GGGGCCCTCTC GGCAATTTTG
 2281 GTGACTCTTG CAAGGTGCTA CTTGGGTGAC GCATCCAAAC TGAGTTCCTC CATCACAGAA
 2341 GGTGTGACCC CCACCCCGCG CCCACGATCA GGAGGCTGGG TCTCCTCCTT CCACCTGCTC
 2401 ACTCCTGGTA GCGCCGGGGG TCGTCCAAGG TTCAAATAGG ACTAGGACCT GTAGTCTGGG
 2461 GTGATCCTGG CTTGACAAGA GGCCTGAGC CTCCCTCTGC AGTTGCGGGC CCGCTTCGGG
 2521 GACGTGTTCA GCCTGCAGCT GGCCTGGAGC CCGGTGGTCG TGCTCAATGG GCTGGCGGCC
 2581 GTGCGCGAGG CGCTGGTGAC CCACGGCCAG GACACCGCCG ACCGCCCCGC TGTGCCCATC
 2641 ACCCAGATCC TGGGTTTTCG GCCGCGTTCC CAAGGCAAGC AGCGGTGGGG ACAGAGACAG
 2701 ATTTCCGTGG GACCCGGGTG GGTGATGACC GTAGTCCGAG CTGGGCAGAG AGGGCGCGGG
 2761 GTCGTGGACA TGAACAGGC CAGCGAGTGG GGACAGCGGG CCAAGAAACC ACCTGCACTA
 2821 GGGAGGTGTG AGCATGGGGA CGAGGGCGGG GCTTGTGAC AGTGGGCGGG GCCACTGCCG
 2881 AGACCTGGCA GGAGCCCAAT GGGTGAGCGT GGCGCATTTT CCAGCTGGAA TCCGTTGTCT
 2941 AAGTGGGGGC GGGGACCGCA CCTGTGCTGT AAGCTCAGTG TGGGTGGCGC GGGGCCCGCG
 3001 GGGTCTTCCC TGAGTGCAAA GGCGGTCAGG GTGGGCAGAG ACAGGTGGG GCAAAGCCTG
 3061 CCCAGCCAA GGGAGCAAGG TGGATGCACA AAGAGTGGGC CCTGTGACCA GCTGGACAGA
 3121 GCCAGGGACT GCGGGAGACC AGGGGGAGCA TAGGGTTGGA GTGGGTGGTG GATGGTGGGG
 3181 CTAATGCCTT CATGGCCACG CGCACGTGCC CGTCCCACCC CCAGGGGTGT TCCTGGCGCG
 3241 CTATGGGCCC GCGTGGCGCG AGCAGAGGCG CTTCTCCGTG TCCACCTTGC GCAACTTGGG
 3301 CCTGGGCAAG AAGTCGCTGG AGCAGTGGGT GACCGAGGAG GCCGCTGCC TTTGTGCCGC
 3361 CTTGCGCAAC CACTCCGGTG GGTGATGGGC AGAAGGGCAC AAAGCGGGAA CTGGGAAGGC
 3421 GGGGGAGCGG GAAGGCGACC CTTACCCACC ATCTCCACCC CCCAGGACGC CCCTTTCGCC
 3481 CCAACGGTCT CTTGGACAAA GCCGTGAGCA ACGTGATCGC CTCCCTCACC TCGGGCGGCC
 3541 GCTTCGAGTA CGACGACCTT CGCTTCCTCA GGCTGCTGGA CCTAGCTCAG GAGGGACTGA
 3601 AGGAGGAGTC GGGCTTTCTG CGCGAGGTGC GGAGCGAGAG ACCGAGGAGT CTCTGCAGGG
 3661 CGAGCTCCCG AGAGGTGCCG GGGCTGACT GGGGCCCTCG AAGAGCAGGA TTTGCATAGA
 3721 TGGGTTTGGG AAAGGACATT CCAGGAGACC CCACTGTAAG AAGGGCTGG AGGAGAGGGG
 3781 GACATCTCAG ACATGCTCGT GGGAGAGGTG TGCCCGGGTC AGGGGGCACC AGGAGAGGCC
 3841 AAGGACTCTG TACCTCCTAT CCACGTGAGA GATTTTCGATT TTAGGTTTCT CCTCTGGGCA
 3901 AGGAGAGAGG GTGGAGGCTG GCACTTGGGG AGGGACTTGG TGAGGTCAAG GGTAAAGGACA

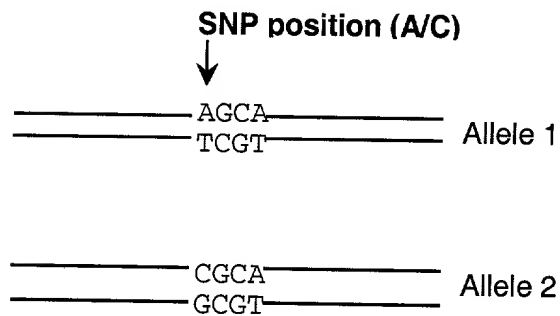
3961	GGCAGGCCCT	GGGTCTACCT	GGAGATGGCT	GGGGCCTGAG	ACTTGTCCAG	GTGAACGCAG
4021	AGCACAGGAG	GGATTGAGAC	CCCCTTCTGT	CTGGTGTAGG	TGCTGAATGC	TGTCCTCCGTC
4081	CTCCTGCATA	TCCAGCGCT	GGCTGGCAAG	GTCTACGCT	TCCAAAAGGC	TTTCTTGACC
4141	CAGCTGGATG	AGCTGCTAAC	TGAGCACAGG	ATGACCTGGG	ACCCAGCCCA	CCCCCCCCGA
4201	GACCTGACTG	AGGCCCTTCCT	GGCAGAGATG	GAGAAGGTGA	GAGTGGCTGC	CACGGTGGGG
4261	GGCAAGGGTG	GTGGGTGAG	CGTCCCAGGA	GGAATGAGGG	GAGGCTGGGC	AAAAGGTTGG
4321	ACCAAGTGCAT	CACCCGGCGA	GCCGCATCTG	GGCTGACAGG	TGCAGAATTG	GAGGTCATTT
4381	GGGGGCTACC	CCGTTCTGTC	CCGAGTATGC	TCTCGGCCCT	GCTCAGGCCA	AGGGGAACCC
4441	TGAGAGCAGC	TTCAATGATG	AGAACCTGCG	CATAGTGGTG	GCTGACCTGT	TCTCTGCCGG
4501	GATGGTGACC	ACCTCGACCA	CGCTGGCCTG	GGGCCTCCTG	CTCATGATCC	TACATCCGGA
4561	TGTGCAGCGT	GAGCCCATCT	GGGAAACAGT	GCAGGGGCCG	AGGGAGGAAG	GGTACAGGCG
4621	GGGGCCCATG	AACCTTGCTG	GGACACCCGG	GGCTCCAAGC	ACAGGCTTGA	CCAGGATCCT
4681	GTAAGCCTGA	CCTCCTCCAA	CATAGGAGGC	AAGAAGGAGT	GTCAGGGCCG	GACCCCTTGG
4741	GTGCTGACCC	ATTGTGGGGA	CGCATGTCTG	TCCAGGCCGT	GTCCAACAGG	AGATCGACGA
4801	CGTGATAGGG	CAGGTGCGGC	GACCAGAGAT	GGGTGACCAG	GCTCACATGC	CCTACACCAC
4861	TGCCGTGATT	CATGAGGTGC	AGCGCTTTGG	GGACATCGTC	CCCCTGGGTG	TGACCCATAT
4921	GACATCCCGT	GACATCGAAG	TACAGGGCTT	CCGCATCCCT	AAGGTAGGCC	TGGCGCCCTC
4981	CTCACCCAG	CTCAGCACCA	GCACCTGGTG	ATAGCCCCAG	CATGGCTACT	GCCAGGTGGG
5041	CCCACTCTAG	GAACCTTGGC	CACCTAGTCC	TCAATGCCAC	CACACTGACT	GTCCCCACTT
5101	GGGTGGGGGG	TCCAGAGTAT	AGGCAGGGCT	GGCCTGTCCA	TCCAGAGCCC	CCGTCTAGTG
5161	GGGAGACAAA	CCAGGACCTG	CCAGAATGTT	GGAGGACCCA	ACGCCTGCAG	GGAGAGGGGG
5221	CAGTGTGGGT	GCCTCTGAGA	GGTGTGACTG	CGCCCTGCTG	TGGGGTCCGA	GAGGGTACTG
5281	TGGAGCTTCT	CGGGCGCAGG	ACTAGTTGAC	AGAGTCCAGC	TGTGTGCCAG	GCAGTGTGTG
5341	TCCCCCGTGT	GTTTGGTGGC	AGGGGTCCCA	GCATCCTAGA	GTCCAGTCCC	CACTCTCACC
5401	CTGCATCTCC	TGCCCAGGGA	ACGACACTCA	TCACCAACCT	GTCATCGGTG	CTGAAGGATG
5461	AGGCCGTCTG	GGAGAAGCCC	TTCCGCTTCT	ACCCCGAACA	CTTCCTGGAT	GCCCAGGGCC
5521	ACTTTGTGAA	GCCGGAGGCC	TTCTCAGGGT	TCTCAGCAGG	TGCCCTGTGG	GAGCCCGGCT
5581	CCCTGTCCCC	TTCCGTGGAG	TCTTGCAGGG	GTATCACCCA	GGAGCCAGGC	TCACTGACGC
5641	CCCTCCCCCT	CCCACAGGCC	GCCGTGCATG	CCTCGGGGAG	CCCCTGGCCC	GCATGGAGCT
5701	CTTCTCTCTC	TTTCACTCCC	TGCTGCAGCA	CTTCAGCTTC	TCGGTGCCCA	CTGGACAGCC
5761	CCGGCCAGC	CACCATGGTG	TCTTTGCTTT	CCTGGTGAGC	CCATCCCCCT	ATGAGCTTTG
5821	TGCTGTGCCC	CGCTAGAATG	GGGTACCTTT	TCCCCAGCCT	GCTCCCTAGC	CAGAGGCTCT
5881	AATGTACAAT	AAAGCAATGT	GGTAGTTCCA	ACTCGGGTCC	CCTGCTCACG	CCCTCGTTGG
5941	GATCATCTCT	CTCAGGGCAA	CCCCACCCCT	GCCTCATTCC	TGCTTACCCC	ACCGCCTGGC
6001	CGCATTTGAG	ACAGGGGTAC	GTTGAGGCTG	AGCAGATGTC	AGTTACCCCT	GCCCATAATC
6061	CCATGTCCCC	CACCTGACCA	ACTCTGACTG	CCCAGATTGG	TGACAAGGAC	TACATTGTCC
6121	TGGCATGTGG	GGAAGGGGCC	AGAATGGGCT	GACTAGAGGT	GTCAAGTCAGC	CTGGATGTG
6181	GTGGAGAGGG	CAGGACTCAG	CCTGGAGGCC	CATATTTTCC	GCCTAACTCA	GCCCCACCCA
6241	CATCAGGGAC	AGCAGTCCCT	CCAGCACCAT	CACAACAGTC	ACCTCCCTTC	ATATATGACA
6301	CCCCAAAACG	GAAGACAAAT	CATGGCGTCA	GGGAGCTATA	TGCCAGGGCT	ACCTACCTCC
6361	CAGGGCTCAG	TCCGGCAGGT	CCAGAACGTT	CCCTGGGAAG	GCCCCATGGA	AGCCCAGGAC
6421	TGAGCCACCA	CCCTCAGCCT	CGTCACCTCA	CCACAGGACT	GGCTACCTCT	CTGGGCCCTC
6481	AGGGATGCTG	CTGTACAGAC	CCCTGACCAG	TGACGAGTTC	GCACTCAGGG	CCAGGCTGGC
6541	GCTGGAGGAG	GACACTTGT	TGGCTCCAAC	CCTAGGTACC	ATCCTCCCAG	TAGGGATCAG
6601	GCAGGGCCCA	CAGGCCCTGCC	CTAGGGACAG	GAGTCAACCT	TGGACCCATA	AGGCACTGGG
6661	GCGGGCAGAG	AAGGAGGAGG	TGGCATGGGC	AGCTGAGAGC	CAGAGACCCT	GACCCTAGTC
6721	CTTGCTCTGC	CATTACCCCG	TGTGACCCCG	GGCCACCCCT	TCCCCACCCCT	TCCCCACCCC
6781	GGGCTTCTGT	TTCTTCTCTG	CAACGAGAAG	GCTGCTTCAC	CTGCCCCGAG	TCCTGTCTTC
6841	CTGCTCTGCC	TTCTGGGGCT	GTGGCCCTTG	CTGGCCTGGA	GCCCCAACCA	AGGGCAGGGA
6901	CTGCTGTCTT	CCACGTCTGT	CCTCACCGAC	ATAATGGGCT	GGGCTGGGCA	CACAGGCAGT
6961	GCCCCAAGAGT	TTCTAATGAG	CATATGATTA	CCTGAGTCCT	GGGCAGACCT	TCTTAGGGAA
7021	CAGCTGGGGA	CAGAGAACCA	CAGACACTCT	GAGGAGCCAC	CCTGAGGCCT	CTTTTGCCAG
7081	AGGACCTTAC	AGCCTCCCTG	GCAGCAGTTC	CGCCAGCATT	TCTGTAAATG	CCCTCATGCC
7141	AGGGTGCGGC	CCGGCTGTCA	GCACGAGAGG	GACGTTGGTC	TGTCCCCTGG	CACCGAGTCA
7201	GTCAGAAGGG	TGGCCAGGGC	CCCCTTGGGC	CCCTCCAGAG	ACAATCCACT	GTGGTCACAC
7261	GGCTCGGTGG	CAGGAAGTGC	TGTTCTGTCA	GCTGTGGGGA	CAGGGAGTGT	GGATGAAGCC
7321	AGGCTGGGTT	TGTCTGAAGA	CGGAGGCCCC	GAAAGGTGGC	AGCCTGGCCT	ATAGCAGCAG
7381	CAACTCTTGG	ATTTATTTGA	AAGATTTTCT	TCACGGTTCT	GAGTCTTGGG	GGTGTTAGAG
7441	GCTCAGAACC	AGTCCAGCCA	GAGCTCTGTC	ATGGGCACGT	AGACCCGGTC	CCAGGGCCTT
7501	TGCTCTTTGC	TGTCCTCAGA	GGCCTCTGCA	AAGTAGAAAC	AGGCAGCCTT	GTGAGTCCCC
7561	TCCTGGGAGC	AACCAACCCCT	CCCTCTGAGA	TGCCCCGGGG	CCAGGTCAGC	TGTGGTGAAA
7621	GGTAGGGATG	CAGCCAGCTC	AGGGAGTGGC	CCAGAGTTCC	TGCCCCACCA	AGGAGGCTCC
7681	CAGGAAGGTC	AAGGCACCTG	ACTCCTGGGC	TGCTTCCCTC	CCCTCCCTTC	CCCAGGTCAG
7741	GAAGGTGGGA	AAGGGCTGGG	TGCTCTGTGA	CCCTGGCAGT	CACTGAGAAG	CAGGGTGGAA
7801	GCAGCCCCCT	GCAGCACGCT	GGGTCACTGG	TCTTACCAGA	TGGATACGCA	GCAACTTCCT
7861	TTTGAACCTT	TTTATTTTCC	TGGCAGGAAG	AAGAGGGATC	CAGCAGTGAG	ATCAGGCAGG
7921	TTCTGTGTTG	CACAGACAGG	GAAACAGGCT	CTGTCCACAC	AAAGTCGGTG	GGGCCAGGAT
7981	GAGGCCAGT	CTGTTACAC	ATGGCTGCTG	CCTCTCAGCT	CTGCACAGAC	GTCTCTGCTC

FIGURE 2

1 GAATTCAAGA CCAGCCTGGA CAACTTGGAA GAACCSGGTC TCTACAAAAA ATACAAAATT
 61 AGCTGGGATT GGGTGCGGTG GCTCATGCCT ATAATCCCAG CACTTTGGGA GCCTGAGGTG
 121 GGTGGATCAC CTGAAGTCAG GAGTTCAAGA CTAGCCTGGC CAACATGGTG AAACCCTATC
 181 TCTACTGAAA ATAYAAAAAG CTAGACGTGG TGGCACACAC CTGTAATCCC AGCTACTTAG
 241 GAGGCTGAGG CAGGAGAATT GCTTGAAGCC TAGAGGTGAA GGTGTAGTG AGCCGAGATT
 301 GCATCATTGC ACAATGGAGG GGAGCCACCA GCCTGGGCAA CAAGAGGAAA TCTCCGTCTC
 361 CAAAAAAAAA AAAAAAAAAA AAAGRATTAG GCTGGGTGGT GCCTGTAGTC CCAGCTACTT
 421 GGGAGGCAGG GGGTCCACTT GATGTCGAGA CTGCAGTGAG CCATGATCCT GCCACTGCAC
 481 TCCGGCCTGG GCAACAGAGT GAGACCCTGT CTAAAGAAAA AAAAAATAAA GCAACATATC
 541 CTGAACAAAG GATCCTCCAT AACGTTCCCA CCAGATTCTT AATCAGAAAC ATGGAGGCCA
 601 GAAAGCAGTG GAGGAGGACR ACCCTCAGGC AGCCCGGGAG GATGTTGTCA CAGGCTGGGG
 661 CAAGGGCCTT CCGGCTACCA ACTGGGAGCT CTGGGAACAG CCCTGTTGCA AACAAGAAGC
 721 CATAGCCCGG CCAGAGCCCA GGAATGTGGG CTGGGCTGGG AGCAGCCTCT GGACAGGAGT
 781 GGTCCCATCC AGGAAACCTC CGGCATGGCT GGGAAAGTGGG GTACTTGGTG CCGGGTCTGT
 841 ATGTGTGTGT GACTGGTGTG TGTGAGAGAG AATGTGTGCY CTAAGTGTCA GTGTGAGTCT
 901 GTGTATGTGT GAATATTGTC TTTGTGTGGG TGATTTCTG CRTGTGTAAT CGTGTCCCTG
 961 CAAGTGTGAA CAAGTGGACA AGTGTCTGGG AGTGGACAAG AGATCTGTGC ACCATCAGGT
 1021 GTGTGCATAG CGTCTGTGCA TGTCAAGAGT GCAAGGTGAA GTGAAGGGAC CAGGCCCATG
 1081 ATGCCACTCA TCATCAGGAG CTCTAAGGCC CCAGGTAAGT GCCAGTGACA GATAAGGGTG
 1141 CTGAAGGTCA CTCTGGAGTG GGCAGGTGGG GGTAAGGAAA GGGCAAGGCC ATGTTCTGGA
 1201 GGAGGGGTG TGACTACATT AGGGTGTATG AGCCTAGCTG GGAGGTGGAT GGCCRGGTCC
 1261 ACTGAAACCC TGGTTATCCC AGAAGGCTTT GCAGGCTTCA GGAGCTTGA GTGGGGAGAG
 1321 GGGGTGACTT CTCCGACCAG GCCCCTCCAC CGGCCTACCC TGGGTAAGGG CCTGGAGCAG
 1381 GAAGCAGGGG CAAGAACCTC TGGAGCAGCC CATACCCGCC CTGGCCTGAC TCTGCCACTG
 1441 GCAGCACAGT CAACACAGCA GGTTCACTCA CAGCAGAGGG CAAAGGCCAT CATCAGCTCC
 1501 CTTTATAAGG GAAGGGTCAC GCGCTCGGTG TGCTGAGAGT GTCCTGCCTG GTCCTCTGTG
 1561 CCTGGTGGGG TGGGGGTGCC AGGTGTGTCC AGAGGAGCCC ATTTGGTAGT GAGGCAGGTA
 1621 TGGGGCTAGA AGCACTGGTG CCCCTGGCCG TGATAGTGGC CATCTTCCTG CTCCTGGTGG

FIGURE 3

One Base Sequencing (OBS) Outline



Add Cy5-ddATP + dTTP, dCTP, dGTP + DNA polymerase

